

Errors Corrected by the STIC Systems Branch

Serial Number: 09/687,483

CRF Processing Date: 10/27/2000
 Edited by: A
 Verified by: A (STIC staff)

ENTERED

RECEIVED

FEB 02 2001

TO 3600 MAIL ROOM

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: _____
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____
- ☒ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: 21
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: _____
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: _____
- ☐ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as _____
- ☐ Inserted mandatory headings, specifically: _____
- ☐ Corrected an obvious error in the response, specifically: _____
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: _____
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted **ending** stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____
- ☐ Other: _____

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

RAW SEQUENCE LISTING DATE: 10/31/2000
 PATENT APPLICATION: US/09/687,483 TIME: 15:27:57

Input Set : A:\Pto.amc
 Output Set: N:\CRF3\10312000\I687483.raw

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3 <110> APPLICANT: Braun et al.
5 <120> TITLE OF INVENTION: METHODS FOR GENERATING DATABASES AND DATABASES FOR IDENTIFYING
6   POLYMORPHIC GENETIC MARKERS
9 <130> FILE REFERENCE: 24736-2033
C--> 11 <140> CURRENT APPLICATION NUMBER: US/09/687,483
12 <141> CURRENT FILING DATE: 2000-10-13
14 <150> PRIOR APPLICATION NUMBER: 60/217,658
15 <151> PRIOR FILING DATE: 2000-07-10
17 <150> PRIOR APPLICATION NUMBER: 60/159,176
18 <151> PRIOR FILING DATE: 1999-10-13
20 <150> PRIOR APPLICATION NUMBER: 60/217,251
21 <151> PRIOR FILING DATE: 2000-07-10
23 <150> PRIOR APPLICATION NUMBER: 09/663,968
24 <151> PRIOR FILING DATE: 2000-09-19
26 <160> NUMBER OF SEQ ID NOS: 118
28 <170> SOFTWARE: FastSEQ for Windows Version 4.0
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31 <211> LENGTH: 361
32 <212> TYPE: DNA
33 <213> ORGANISM: Homo Sapien
35 <400> SEQUENCE: 1
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37 agcaatggat gatttgatgc tgtccccgga cgatattgaa caatggttca ctgaagaccc      120
38 aggtccagat gaagctccca gaatgccaga ggctgctccc cgcgtggccc ctgcaccagc      180
39 agctcctaca ccggcgggccc ctgcaccagc cccctcctgg ccctgtcat cttctgtccc      240
40 ttcccagaaa acctaccagg gcagctacgg ttccggtctg ggcttcttgc attctgggac      300
41 agccaagtct gtgacttyca cgttcagttg ccctgagggg ctggcttcca tgagacttca      360
42 a
44 <210> SEQ ID NO: 2
45 <211> LENGTH: 44
46 <212> TYPE: DNA
47 <213> ORGANISM: Artificial Sequence
49 <220> FEATURE:
50 <223> OTHER INFORMATION: Oligonucleotide Primer
52 <400> SEQUENCE: 2
53 cccagtcacg acgttgtaaa acgctgagga cctggctctc tgac      44
55 <210> SEQ ID NO: 3
56 <211> LENGTH: 42
57 <212> TYPE: DNA
58 <213> ORGANISM: Artificial Sequence
60 <220> FEATURE:
61 <223> OTHER INFORMATION: Oligonucleotide Primer
63 <400> SEQUENCE: 3
64 agcggataac aatttcacac aggttgaagt ctcatggaag cc      42
66 <210> SEQ ID NO: 4
67 <211> LENGTH: 17
68 <212> TYPE: DNA

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RAW SEQUENCE LISTING DATE: 10/31/2000
 PATENT APPLICATION: US/09/687,483 TIME: 15:27:57

Input Set : A:\Pto.amc
 Output Set: N:\CRF3\10312000\I687483.raw

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69 <213> ORGANISM: Artificial Sequence
71 <220> FEATURE:
72 <223> OTHER INFORMATION: Probe
74 <400> SEQUENCE: 4
75 gccagaggct gctccccc 17
77 <210> SEQ ID NO: 5
78 <211> LENGTH: 17
79 <212> TYPE: DNA
80 <213> ORGANISM: Artificial Sequence
82 <220> FEATURE:
83 <223> OTHER INFORMATION: Probe
85 <400> SEQUENCE: 5
86 gccagaggct gctccccc 17
88 <210> SEQ ID NO: 6
89 <211> LENGTH: 19
90 <212> TYPE: DNA
91 <213> ORGANISM: Artificial Sequence
93 <220> FEATURE:
94 <223> OTHER INFORMATION: Probe
96 <400> SEQUENCE: 6
97 gccagaggct gctcccgc 19
99 <210> SEQ ID NO: 7
100 <211> LENGTH: 18
101 <212> TYPE: DNA
102 <213> ORGANISM: Artificial Sequence
104 <220> FEATURE:
105 <223> OTHER INFORMATION: Probe
107 <400> SEQUENCE: 7
108 gccagaggct gctccccc 18
110 <210> SEQ ID NO: 8
111 <211> LENGTH: 161
112 <212> TYPE: DNA
113 <213> ORGANISM: Homo Sapien
115 <400> SEQUENCE: 8
116 gtccgtcaga acccatgcgg cagcaaggcc tgcgcgcgc tcttcggccc agtggacagc 60
117 gaggagctga gccgcgactg tgatgcgcta atggcgggct gcatccagga ggcccgtgag 120
118 cgatgggaact tcgactttgt caccgagaca ccaactggagg g 161
120 <210> SEQ ID NO: 9
121 <211> LENGTH: 43
122 <212> TYPE: DNA
123 <213> ORGANISM: Artificial Sequence
125 <220> FEATURE:
126 <223> OTHER INFORMATION: Oligonucleotide Primer
128 <400> SEQUENCE: 9
129 cccagtcacg acgttgtaaa acggtccgtc agaaccatg cgg 43
131 <210> SEQ ID NO: 10
132 <211> LENGTH: 44
133 <212> TYPE: DNA
134 <213> ORGANISM: Artificial Sequence

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RAW SEQUENCE LISTING DATE: 10/31/2000
 PATENT APPLICATION: US/09/687,483 TIME: 15:27:57

Input Set : A:\Pto.amc
 Output Set: N:\CRF3\10312000\I687483.raw

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136 <220> FEATURE:
137 <223> OTHER INFORMATION: Oligonucleotide Primer
139 <400> SEQUENCE: 10
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142 <210> SEQ ID NO: 11
143 <211> LENGTH: 15
144 <212> TYPE: DNA
145 <213> ORGANISM: Artificial Sequence
147 <220> FEATURE:
148 <223> OTHER INFORMATION: Oligonucleotide Primer
150 <400> SEQUENCE: 11
151 cagcgagcag ctgag      15
153 <210> SEQ ID NO: 12
154 <211> LENGTH: 15
155 <212> TYPE: DNA
156 <213> ORGANISM: Artificial Sequence
158 <220> FEATURE:
159 <223> OTHER INFORMATION: Probe
161 <400> SEQUENCE: 12
162 cagcgagcag ctgag      15
164 <210> SEQ ID NO: 13
165 <211> LENGTH: 16
166 <212> TYPE: DNA
167 <213> ORGANISM: Artificial Sequence
169 <220> FEATURE:
170 <223> OTHER INFORMATION: Probe
172 <400> SEQUENCE: 13
173 cagcgagcag ctgagc      16
175 <210> SEQ ID NO: 14
176 <211> LENGTH: 17
177 <212> TYPE: DNA
178 <213> ORGANISM: Artificial Sequence
180 <220> FEATURE:
181 <223> OTHER INFORMATION: Probe
183 <400> SEQUENCE: 14
184 cagcgagcag ctgagc      17
186 <210> SEQ ID NO: 15
187 <211> LENGTH: 205
188 <212> TYPE: DNA
189 <213> ORGANISM: Homo Sapien
191 <400> SEQUENCE: 15
192 gcgctccatt catctcttca tgcactctct gttgaatgaa gaaaatccaa gtaaggccta      60
193 caggtgcagt tccaagggaag cctttgagaa agggctctgc ttgagttgta gaaagaaccg      120
194 ctgcaacaat ctgggctatg agatcaataa agtcagagcc aaaagaagca gcaaaatgta      180
195 cctgaagact cgttctcaga tgccc      205
197 <210> SEQ ID NO: 16
198 <211> LENGTH: 42
199 <212> TYPE: DNA
200 <213> ORGANISM: Artificial Sequence

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RAW SEQUENCE LISTING DATE: 10/31/2000
 PATENT APPLICATION: US/09/687,483 TIME: 15:27:57

Input Set : A:\Pto.amc
 Output Set: N:\CRF3\10312000\I687483.raw

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202 <220> FEATURE:
203 <223> OTHER INFORMATION: Oligonucleotide Primers
205 <400> SEQUENCE: 16
206 cccagtcacg acgttgtaaa acggcgctcc attcatctct tc      42
208 <210> SEQ ID NO: 17
209 <211> LENGTH: 42
210 <212> TYPE: DNA
211 <213> ORGANISM: Artificial Sequence
213 <220> FEATURE:
214 <223> OTHER INFORMATION: Oligonucleotide Primer
216 <400> SEQUENCE: 17
217 agcggataac aatttcacac agggggcatc tgagaacgag tc      42
219 <210> SEQ ID NO: 18
220 <211> LENGTH: 20
221 <212> TYPE: DNA
222 <213> ORGANISM: Artificial Sequence
224 <220> FEATURE:
225 <223> OTHER INFORMATION: Oligonucleotide Primer
227 <400> SEQUENCE: 18
228 caatctgggc tatgagatca      20
230 <210> SEQ ID NO: 19
231 <211> LENGTH: 20
232 <212> TYPE: DNA
233 <213> ORGANISM: Artificial Sequence
235 <220> FEATURE:
236 <223> OTHER INFORMATION: Probe
238 <400> SEQUENCE: 19
239 caatctgggc tatgagatca      20
241 <210> SEQ ID NO: 20
242 <211> LENGTH: 21
243 <212> TYPE: DNA
244 <213> ORGANISM: Artificial Sequence
246 <220> FEATURE:
247 <223> OTHER INFORMATION: Probe
249 <400> SEQUENCE: 20
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252 <210> SEQ ID NO: 21
253 <211> LENGTH: 22
254 <212> TYPE: DNA
255 <213> ORGANISM: Artificial Sequence
257 <220> FEATURE:
258 <223> OTHER INFORMATION: Probe
260 <400> SEQUENCE: 21
261 caatctgggc tatgagatca gt      22
263 <210> SEQ ID NO: 22
264 <211> LENGTH: 60
265 <212> TYPE: DNA
266 <213> ORGANISM: Homo Sapien
268 <220> FEATURE:

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RAW SEQUENCE LISTING DATE: 10/31/2000
 PATENT APPLICATION: US/09/687,483 TIME: 15:27:57

Input Set : A:\Pto.amc
 Output Set: N:\CRF3\10312000\I687483.raw

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269 <223> OTHER INFORMATION: Probe
271 <400> SEQUENCE: 22
272 qtgccggcta ctccggatggc agcaaggact cctgcaaggg ggacagtgga ggcacacatg      60
274 <210> SEQ ID NO: 23
275 <211> LENGTH: 60
276 <212> TYPE: DNA
277 <213> ORGANISM: Homo sapien
279 <400> SEQUENCE: 23
280 ccaccacta ccggggcagc tggtagctga cgggcatcgt cagctggggc cagggtctgcg      60
282 <210> SEQ ID NO: 24
283 <211> LENGTH: 42
284 <212> TYPE: DNA
285 <213> ORGANISM: Artificial Sequence
287 <220> FEATURE:
288 <223> OTHER INFORMATION: Oligonucleotide primer
290 <400> SEQUENCE: 24
291 ccagtcacg acgttgtaaa acgatggcag caaggactcc tg                          42
293 <210> SEQ ID NO: 25
294 <211> LENGTH: 18
295 <212> TYPE: DNA
296 <213> ORGANISM: Artificial Sequence
298 <220> FEATURE:
299 <223> OTHER INFORMATION: Oligonucleotide primer
301 <400> SEQUENCE: 25
302 cacatgccac ccactacc                                          18
304 <210> SEQ ID NO: 26
305 <211> LENGTH: 43
306 <212> TYPE: DNA
307 <213> ORGANISM: Artificial Sequence
309 <220> FEATURE:
310 <223> OTHER INFORMATION: Oligonucleotide primer
312 <400> SEQUENCE: 26
313 agcggataac aatttcacac aggtgacgat gcccgtcagg tac                      43
315 <210> SEQ ID NO: 27
316 <211> LENGTH: 15
317 <212> TYPE: DNA
318 <213> ORGANISM: Artificial Sequence
320 <220> FEATURE:
321 <223> OTHER INFORMATION: Probe
323 <400> SEQUENCE: 27
324 atgccacca ctacc                                          15
326 <210> SEQ ID NO: 28
327 <211> LENGTH: 19
328 <212> TYPE: DNA
329 <213> ORGANISM: Artificial Sequence
331 <220> FEATURE:
332 <223> OTHER INFORMATION: Probe
334 <400> SEQUENCE: 28
335 cacatgccac ccactaccg                                          19

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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/687,483

DATE: 10/31/2000

TIME: 15:27:58

Input Set : A:\Pto.amc

Output Set: N:\CRF3\10312000\I687483.raw

L:11 M:270 C: Current Application Number differs, Replaced Current Application Number
L:7310 M:283 W: Missing Blank Line separator, <210> field identifier

OIPE

RAW SEQUENCE LISTING DATE: 10/27/2000
 PATENT APPLICATION: US/09/687,483 TIME: 08:33:09

Input Set : A:\2033seq.001
 Output Set: N:\CRF3\10272000\I687483.raw

Does Not Comply
 Corrected Diskette Needed

3 <110> APPLICANT: Braun et al.
 5 <120> TITLE OF INVENTION: METHODS FOR GENERATING DATABASES AND DATABASES FOR IDENTIFYING
 6 POLYMORPHIC GENETIC MARKERS
 9 <130> FILE REFERENCE: 24736-2033
 C--> 11 <140> CURRENT APPLICATION NUMBER: US/09/687,483
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 17 <150> PRIOR APPLICATION NUMBER: 60/159,176
 18 <151> PRIOR FILING DATE: 1999-10-13
 20 <150> PRIOR APPLICATION NUMBER: 60/217,251
 21 <151> PRIOR FILING DATE: 2000-07-10
 23 <150> PRIOR APPLICATION NUMBER: 09/663,968
 24 <151> PRIOR FILING DATE: 2000-09-19
 26 <160> NUMBER OF SEQ ID NOS: 118
 28 <170> SOFTWARE: FastSEQ for Windows Version 4.0

ERRORED SEQUENCES

252 <210> SEQ ID NO: 21
 253 <211> LENGTH: 22
 254 <212> TYPE: DNA
 255 <213> ORGANISM: Artificial Sequence
 257 <220> FEATURE:
 258 <223> OTHER INFORMATION: Probe
 260 <400> SEQUENCE: 21
 E--> 261 caatctgggc tatgagatca gt

20 22

VERIFICATION SUMMARY DATE: 10/27/2000
PATENT APPLICATION: US/09/687,483 TIME: 08:33:12

Input Set : A:\2033seq.001
Output Set: N:\CRF3\10272000\I687483.raw

L:11 M:270 C: Current Application Number differs, Replaced Current Application Number
L:261 M:254 E: No. of Bases conflict, LENGTH:Input:20 Counted:22 SEQ:21